FIG. 3A

FIG. 3B

666 000 019 810

FIG. 3C

GCT CGA Ala

FIG. 3D

FIG. 3E

1350	ATT 6GT TAA CCA 11e Gly>	1440 AAA 6CC TIT CG6 Lys Ala>	AGA ACG TCT TGC Arg Thr>
	AAC TTG ASD	ATT TAA TIe	AAG Phe C
1340	GAT AAC CTA TTG ASD ASD	1430 AAA CAC TIT 616 Lys H1s	1520 T ATC TTA A TAG AAT S 11e Leu
•	A TGG	A 660 T 666 U 619	G CAT C GTA n His
1330	CAG CAA GTC GTT GID GID	1420 6AA 6AA CTT CTT 61u 61u	1510 ACT CAG TGA GTC Thr GIn
•	CAG 610 610	AAAA Phe	AAA TTT Lys
	ACT TGA Thr	000 000 0 1 1 2 0 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	600 000 000 000 000 000 000 000
1320	GCT ATT CGA TAA Ala Ile IDE-C	1410 616 6AT 7 CAC CTA 7 VAI ASP 11DE_CC	1500 6 66T AAA C CCA TTT F 61y Lys TIDE_C
	C ATC ATC TAG TAG PEPT	C TAT G ATA a Tyr PEPT	TCC N AGG
1310	CGT GTC GCA CAG Arg Val (OTOXIN F	AAA GCC TTT CGG Lys Ala KOTOXIN B	1490 AGT AAT TCA TTA Ser ASA KOTOXIN
•	CTTT CTTT LEU	661 • CCA 61V LEUJ	GTG CAC CAC LEUR
1300	CAG GCA GTC CGT GIN Ala COMBINANT	1390 CCTT AGT GAA TCA I Leu Ser COMBINANT	1480 T ATT GAT A TAA CTA E IJE ASD COMBINANT
•	C AAT U Leu L-RECC	A GTC T CAG S Val	T ATT Y INE
1290	AAA GAG TTT CTC Lys Glu	GAA AAA CTT TTT GIU LYS	AAC GGT TTG CCA ASN GIY
, •	AAT TAG AAT TTG Leu Asn	TTA GGT AAT CCA Leu Gly	TCG 6CA AGC CGT Ser Ala
1280	AAT ATG AAA TTC TTA CTG AAC TTA AAC AAA TTA TAA TTA TTA TTG TTT ASG AAT GAC TTG AAT TTG TTT ASM Met Lys Phe Leu Leu Asm Leu Asm Lys Locardon Comments of the C	1370 1. T AGC CGT TTA GGT A TCG GCA AAT CCA e Ser Arg Leu Gly 	1450 1460 1470 TTA GTA CAG TTG GAT TCG GCA AAC AAT CAT GTC AAC CTA AGC CGT TTG Leu Val Gln Leu Asp Ser Ala Asn
•	VAT G	ATT P	CA6 GTC GIN
1270	Y TTC 1 F AAG / S Phe L	1360 GAT TTA GCT GGT ATT A CTA AAT CGA CCA TAA T ASD Leu Ala GIY Ile S	1450 TA GTA AT CAT eu Val
12	G AA/ C TT1 t Lys	15 A 667 U A16	
•	A TAI	A AA AA Le	GAT AAA CTA TTT ASP LYS
	AA TT AS	20.8 I	304 .

FIG. 3F

FIG. 3G

FIG. 3H

2170 2230 2240 2250 2250 2250 2250 2250 2250 225	CAC GGT GGC AAG GGC GAT GAT ATT TTC GTT CAC CGT AAA GGC GAT GGT AAT GAT ATT ATT ACC GAT TCT GAC GGC AAT GAT AAA TTA TCA GGT CCA CGC TTA CTA TTA TAA TAA TAA TAAT AGT TTA AAT AGT TTA AAT AGT CCA CCA TTC CCA CCA TTC CCA TTA CTA TAA TA	2350 2410 2420 2430 2430 2390 2490 2400 2410 2420 2430 2430 2430 2400 2410 2420 2430 2430 2430 2430 2430 2430 243	CAA AAC TGG TTC CGA GAG GCT GAT TTT GCT AAA GAA GTG CCT AAT TAT AAA GCA ACT AAA GAT GAG AAA ATC GAA GAA ATC AAG GTC CAA GGT CA
GAC CGC TTA TTT GGT GGT AAA CTG GCG AAT AAA CCA CCA TTA ASD Arg Leu Phe Gly Gly Lys	2260	2350	2440
	CAC GGT GGC AAG GGC	TTC TCT GAT TCG AAC	CAA AAC TGG TTC CG/
	GTG CCA CCG TTC CCG	AAG AGA CTA AGC TTG	GTT TTG ACC AAG GC/
	His Gly Gly Lys Gly	Phe Ser Asp Ser Asr	GIn Asn Trp Phe Arg

FIG. 33